

E. Coli

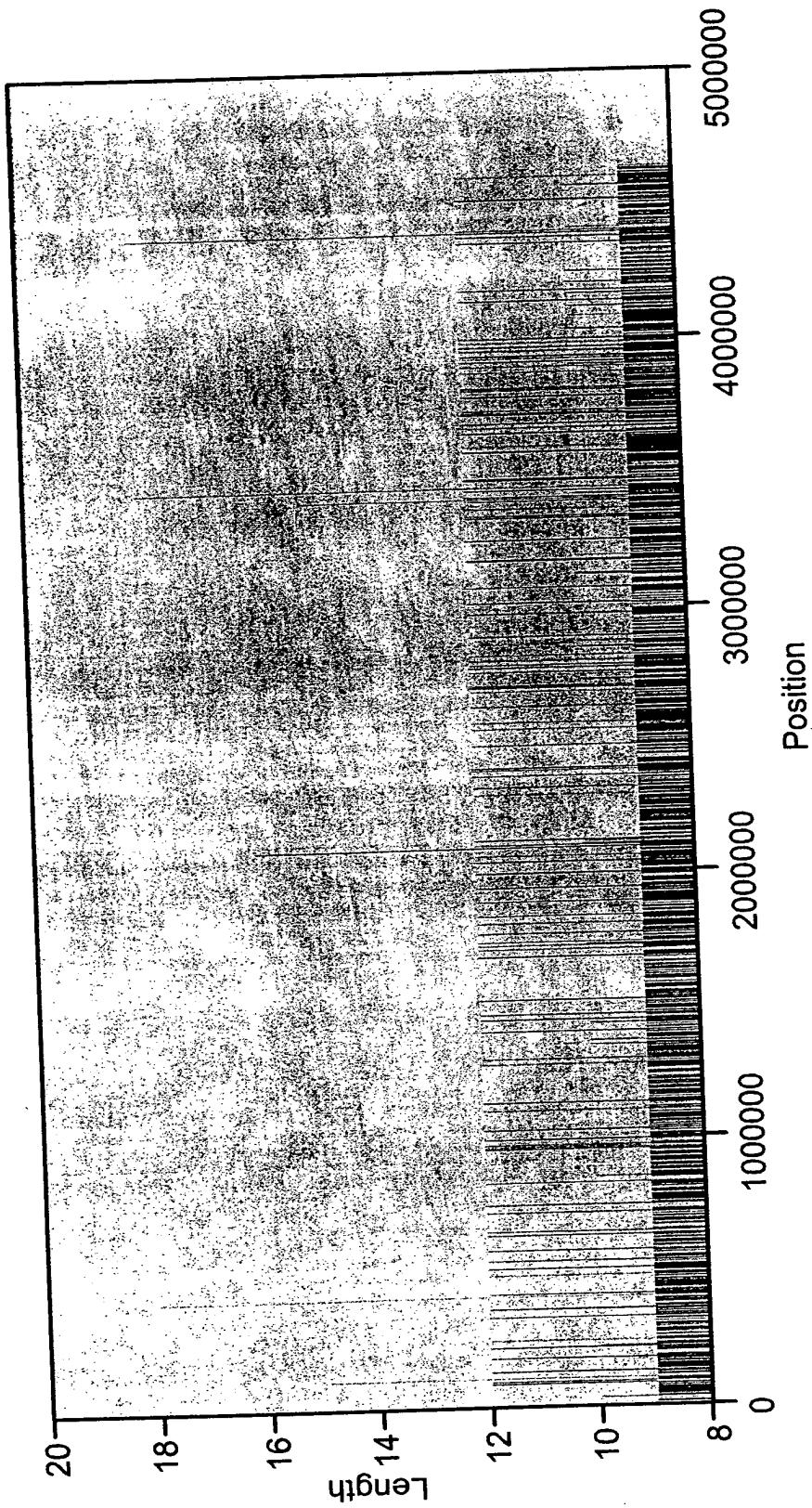


Fig. 1a

*Bacillus Subtilis*

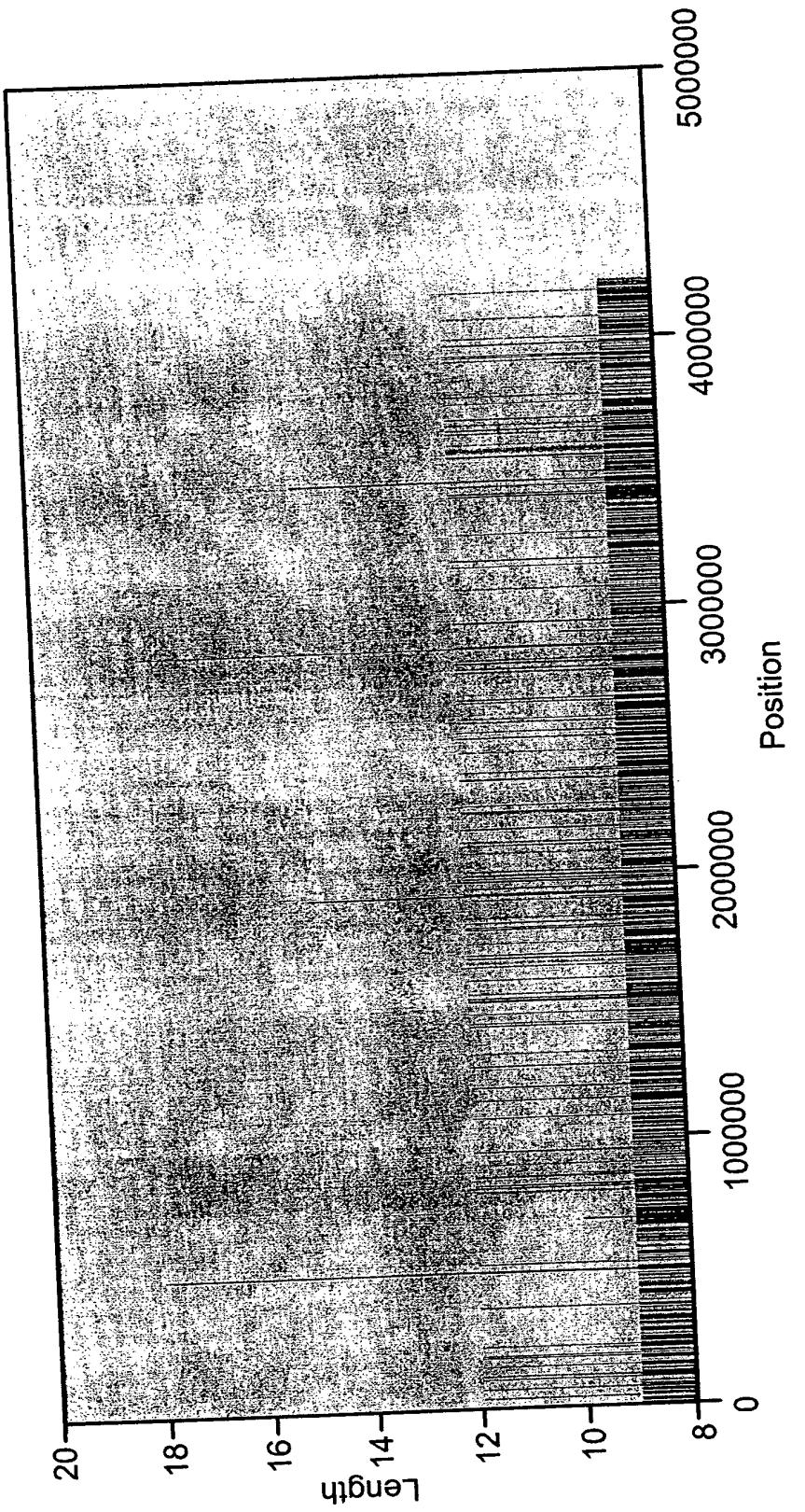


Fig. 1b

*Archaeoglobus fulgidus*

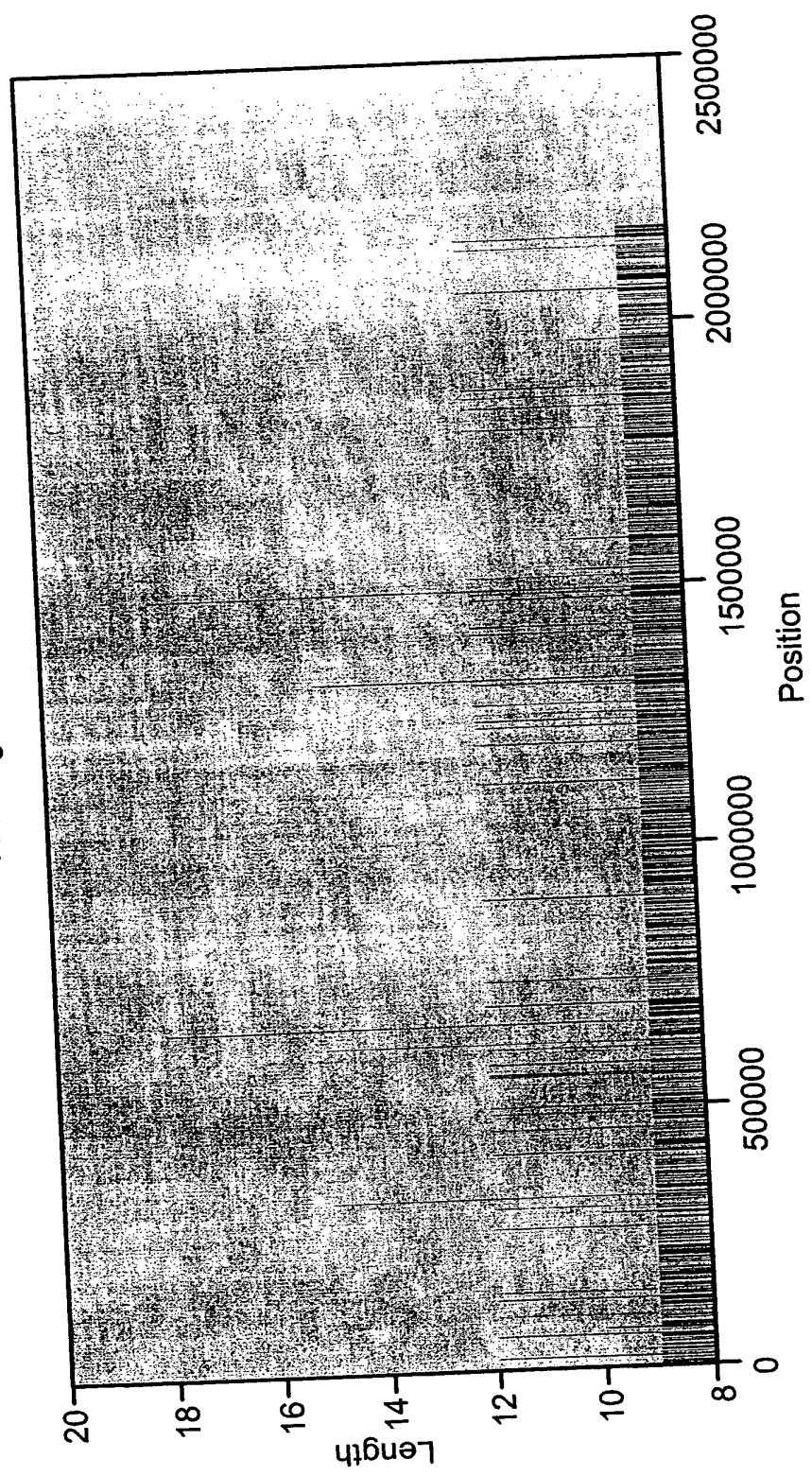


Fig. 1c

Yeast Chromosome 7

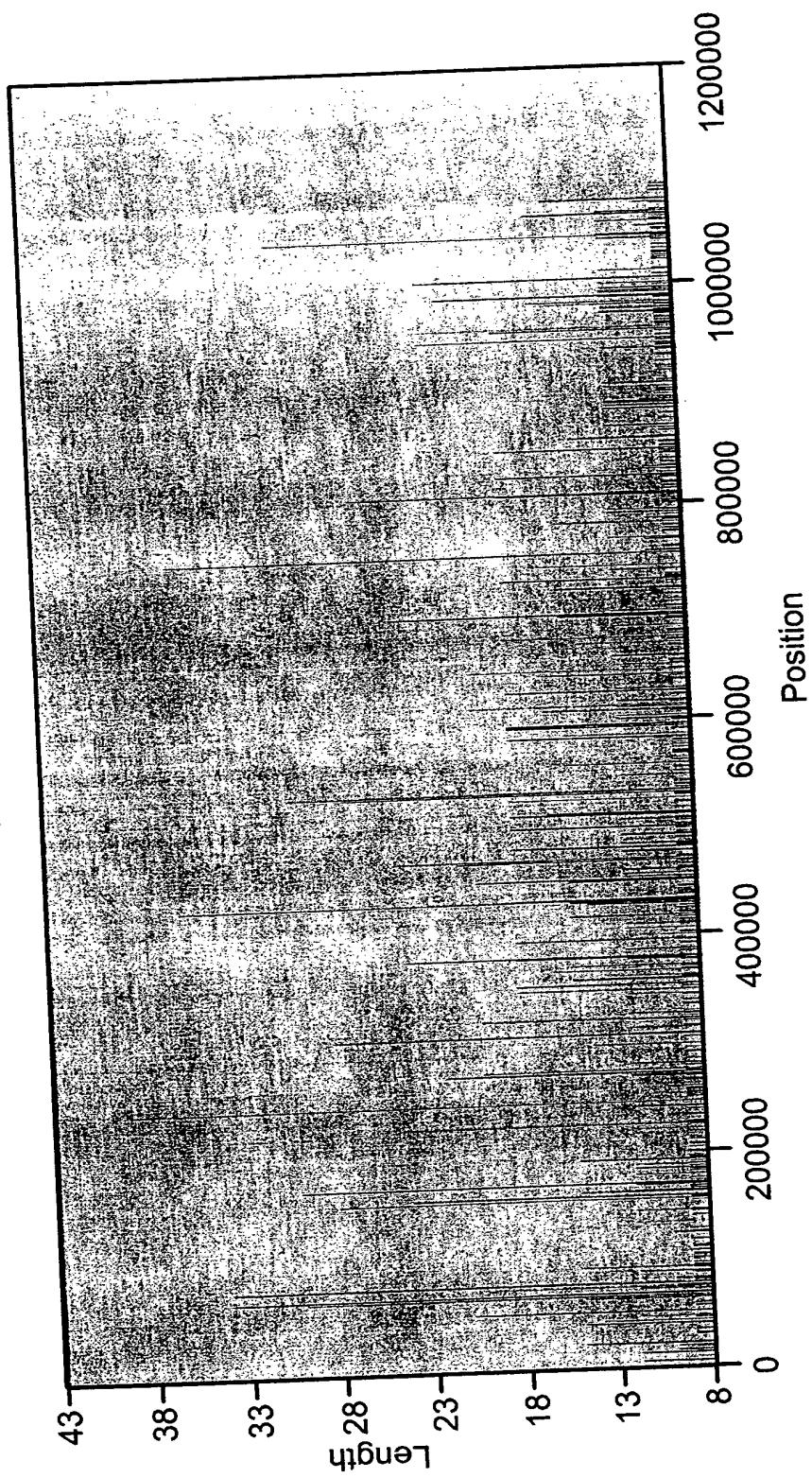


Fig. 1d

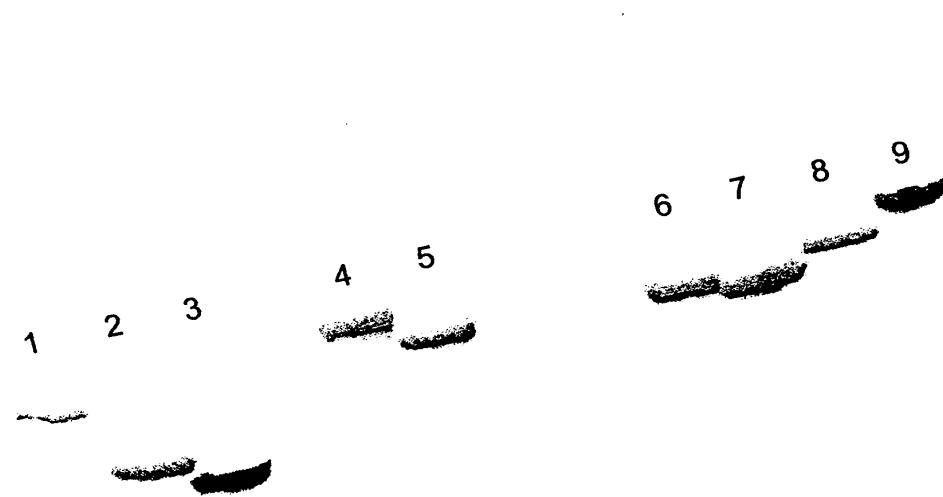


Fig. 2

		50
Ec K12, DH5 $\alpha$	1	---GTTATGT CTTATCCCAC GGTATTTAAT ATGGTTCAATT AGGATGTTTA 25*
Ec Bsr9b		-TTGTTATGT CTTATCCCAC GGTATTTAAT ATAGTTCAATT TGGATGTTCA 26*
Ec Bsr9c		TTTGTATGT CTTATCCCAC GGTATTTAAT ATAGTTCAATT TGGATGTTCA 27*
Ec ETEC		-TCTATGTT CTTATCNCCAC GGTNTTTAAT ATGGTTCAATT AGGATGTTTA 28*
Consensus		-----T-T--- -T---CCCAC GGTATTTAAT AT-GTTCAATT -GGATGTT-A 29*
		100
Ec K12, DH5 $\alpha$	51	TTTCTTGATT TTGCATATGA GTATATTA.. CCCCCCCCCC AAAAAAATAA
Ec Bsr9b		TTTCTTTATT TTGCATATGA GTATATTA.. .CCCCTT CAAAAAATAA
Ec Bsr9c		TTTCTTTATT TTGCATATGA GTATATTA.. .CCCCTT CAAAAAATAA
Ec ETEC		TTTCTTGATT TTGCATATGA GTATATTACC CCCCCCCCCC AAAAAAATAA
Consensus		TTTCTT-ATT TTGCATATGA GTATATTA-- ---CCCCT- -AAAAAATAA
		150
Ec K12, DH5 $\alpha$	101	ATTAATTAAA ATGATGGCTT ATATAAAAATA AAATTTAAAG CAAGGAATCT
Ec Bsr9b		ATTAATTAAA ACGATTGCTT ATATAAAAACA AAATTTAAAG CAAGGAATCT
Ec Bsr9c		ATTAATTAAA ACGATTGCTT ATATAAAAACA AAATTTAAAG CAAGGAATCT
Ec ETEC		ATTAATTAAA ATGATGGCTT ATATNAAAATA AAATTTAAAG CAAGGANTCT
Consensus		ATTAATTAAA A-GAT-GCTT ATATAAAA-A AAATTTAAAG CAAGGAATCT
		200
Ec K12, DH5 $\alpha$	151	CAATGGATGT TAAACAAAAT GAGATTTGT GAAAGCAATA AATTATTGAC
Ec Bsr9b		CAATGGATGT TAAACAAAAT GAGATTTAGT GAAAACAATA AATTATTCAC
Ec Bsr9c		CAATGGATGT TAAACAAAAT GAGATTTAGT GAAAACAATA AATTATTCAC
Ec ETEC		CAATGGATGT TAAACANAAT GAGATTTGT GAANGCNATN NATTATTGNC
Consensus		CAATGGATGT TAAACAAAAT GAGATTT-GT GAAA-CAATA AATTATT-AC
		250
Ec K12, DH5 $\alpha$	201	TTCGTTTTAG ATTTGTTAG CTATAATGTT ATACATTCAA ATGACTGAAC
Ec Bsr9b		TTCGTTTTAG ATTTGTTAG CTATAATGTT ATACATTCAA ATGACTGAAC
Ec Bsr9c		TTCGTTTTAG ATTTGTTAG CTATAATGTT ATACATTCAA ATGACTGAAC
Ec ETEC		TTCGTTGTAN ATTTGCTNAG CTATAATGTT ATNCATTCAA ATGACTGAAC
Consensus		TTCGTT-TAG ATTTG-TTAG CTATAATGTT ATACATTCAA ATGACTGAAC
		264
Ec K12 DH5 $\alpha$	251	ATCCTGTAAT TAAA
Ec Bsr9b		ATCCTGTATT TAA-
Ec Bsr9c		ATCCTGTAAT TAA-
Ec ETEC		ATCCTGTNNT TANA
Consensus		ATCCTGTAAT TAA-

\* SEQ ID NO

Fig. 3a

	1	50				
<i>Ec K12, DH5<math>\alpha</math></i>	TTTNCCCGA	AAAAAAATAGG	AAAGGGGGGG	GGGCTAATCG	GCAGGGAAAGG	30*
<i>Ec K12, w3110</i>	TNTTNNNCGG	AAAAAAATNG	AAAGGGGGGG	GGGCTAATCG	GCAGGGAAAGG	31*
<i>Ec Bsr9c</i>	--TTTNCCCGA	AAAAAAATNG	AAA..GGGGG	GGGCTAATCG	GCAGGGAAAGG	32*
<i>Ec (wt) 1</i>	--TNTNCGGA	AAAAAAANAGG	AAAGGGGGGG	GGGCTAATCG	GCAGGGAAAGG	33*
<i>Ec (wt) 54</i>	-----NCG	AAAAAAATG	AAA.GGGGGG	GGGCTAATCG	GCAGGGAAAGG	34*
<i>Ec (wt) 68</i>	-----CG	AAAAAAATG	AAA.GGGGGG	GGGCTAATCG	GCAGGGAAAGG	35*
Consensus	-----	-AAAAAA--G	AAA--GGGGG	GGGCTAATCG	GCAGGGAAAGG	36*
	51	100				
<i>Ec K12, DH5<math>\alpha</math></i>	CCGCCCCCGA	TAGCGGGCGG	CANAAGGAAT	CANAATTTC	AGGTCAGACG	
<i>Ec K12, w3110</i>	CCGCCCCCGA	TAGCGGGCGG	CAGAAGGAAT	CAGAATTTC	AGGTCAGACG	
<i>Ec Bsr9c</i>	CCGCCCCCGA	TAGCGGGCGG	CAGAAGGAAT	CAGAATTTC	AGGTCAGATG	
<i>Ec (wt) 1</i>	CCGCCCCCGA	TAGCGGGCGG	CAGAAGGAAT	CAGAATTTC	AGGTCAGACG	
<i>Ec (wt) 54</i>	CCGCCCCCGA	TAGCGGGCGG	CAGAAGGAAT	CAGAATTTC	AGGTCAGATG	
<i>Ec (wt) 68</i>	CCGCCCCCGA	TAGCGGGCGG	CAGAAGGAAT	CAGAATTTC	AGGTCAGATG	
Consensus	CCGCCCCCGA	TAGCGGGCGG	CAGAAGGAAT	CAGAATTTC	AGGTCAGA-G	
	101	150				
<i>Ec K12, DH5<math>\alpha</math></i>	GGCTGCAAGT	TGCAGACCGT	TAAAATCATC	GGNNNGGGTG	TCGTACCAACA	
<i>Ec K12, w3110</i>	GGCTGCAAGT	TGCAGACCGT	TAAAATCATC	GGTTGGGGTG	TCGTACCAACA	
<i>Ec Bsr9c</i>	GGCTGCAAGT	TGCAGACCGT	TATAATCATC	GGTTGGGGTG	TCGTACCAACA	
<i>Ec (wt) 1</i>	GGCTGCAAGT	TGCAGACCGT	TAAAATCATC	GGTTGGGGTG	TCGTACCAACA	
<i>Ec (wt) 54</i>	GGCTGCAAGT	TGCAGACCGT	TATAATCATC	GGTTGGGGTG	TCGTACCAACA	
<i>Ec (wt) 68</i>	GGCTGCAAGT	TGCAGACCGT	TATAATCATC	GGTTGGGGTG	TCGTACCAACA	
Consensus	GGCTGCAAGT	TGCAGACCGT	TA-AATCATC	GGTTGGGGTG	TCGTACCAACA	
	151	180				
<i>Ec K12, DH5<math>\alpha</math></i>	CTTTACCTGC	CGTCAGCCCG	AGATTTAA-GTT	-G		
<i>Ec K12, w3110</i>	CTTTACCTGC	CGTCAGCCCG	AGATTTAA-GTT	-G		
<i>Ec Bsr9c</i>	CTTTACCTGC	CGTCAGCCCG	AGATTTAA-GTT	-G		
<i>Ec (wt) 1</i>	CTTTACCTGC	CGTCAGCCCG	AGATTTAAAGTT	TGG		
<i>Ec (wt) 54</i>	CTTTACCTGC	CGTCAGCCCG	AGAT-AAAGTT	TGG		
<i>Ec (wt) 68</i>	CTTTACCTGC	CGTCAGCCCG	AGAT-AAAGTT	TGG		
Consensus	CTTTACCTGC	CGTCAGCCCG	AGAT-AA-GTT	-G		

\* SEQ ID NO

Fig. 3b

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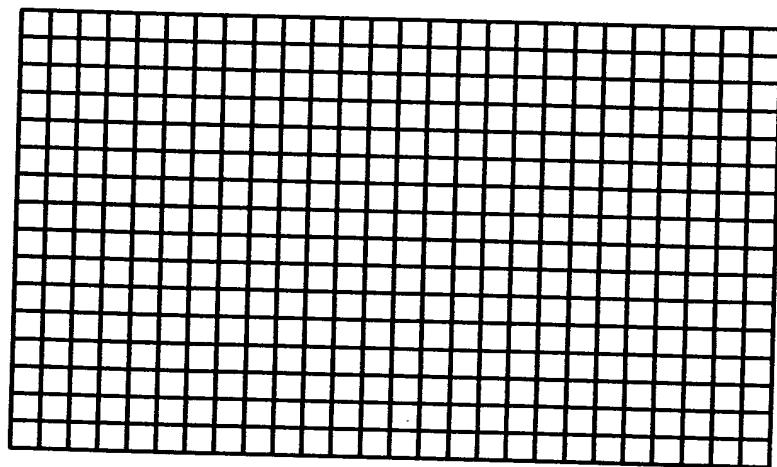


Fig. 4